IN SEARCH OF: A MEMBRANE SIALOGLYCOPROTEIN CELL GROWTH INHIBITOR

bу

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SECTION I
ACKNOWLEDGEMENTS

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SECTION II
HISTORICAL REVIEW

HISTORICAL REVIEW

Density-dependent Cell Growth

Cell-cell interactions are involved in the regulation of cell growth. The study of such cellular interactions and cell growth control was simplified by the development of in vitro cultivation. Eukaryotic cells were first successfully grown in culture in the early 1900s by Harrison (1907) and Carrel (1912), and these earliest studies of cultured cells included simple observations of cellular behavior in this unnatural environment. Abercrombie and Heaysman (1954) described the formation of "monolayers" of cells grown in hanging drop slides and demonstrated that the cellular arrangement into monolayers resulted from "contact inhibition." Contact inhibition referred to the phenomenon whereby cellular movement was restricted through cell-cell contact. Since that initial observation, numerous researchers have explored the causes of such cell behavior.

Further investigations suggested that not only was cell motility diminished, but also cell proliferation was reduced upon cell-cell contact. Later, these two events

were seen to be separate; Stoker and Rubin (1967) proposed terminology to distinguish between contact inhibition, meaning restricted movement of cells, and density-dependent inhibition, meaning restricted growth of cells. Subsequently, however, these terms have been used inter-changeably in reference to cell growth control.

A loss of contact or density-dependent inhibition of cell growth was correlated to tumorigenicity of cell populations. Aaronson and Todaro (1968) found that cell lines which grew efficiently at high saturation densities in vitro produced tumors in vivo more frequently than cell lines which demonstrated contact inhibition in culture. About the same time, Holley and Kiernan (1968) demonstrated that contact inhibition of cells in culture could be overcome by serum factors. They found that the final cell density was directly proportional to the concentration of serum in the medium. In other words, the more serum (thus, serum factors) present, the greater the number of cells produced in a given time and space. Such growth-promoting molecules have been the object of many investigative pursuits.

Positive Growth Regulators

Enhancement of cellular proliferation has been attributed to many agents including known transforming substances (such as viruses and phorbol esters), serum factors and cell-associated molecules. The numerous transforming substances are beyond the scope of this review and the serum components have not yet been completely identified; therefore, only those agents isolated from cells will be discussed here.

The majority of cell-derived molecules have been described as polypeptide growth factors. These polypeptide growth factors are similar to hormones in that they are relatively small polypeptides and that they are released and received by cells. Several such molecules have been isolated, purified and characterized in detail; while others have been purified and/or characterized to a lesser extent.

One of the first of these molecules to be isolated and carefully studied was epidermal growth factor (EGF). EGF activity was initially discovered about thirty years ago in an extract of mouse submaxillary glands by Cohen (1962), while he was studying nerve growth factor (NGF). Later, Cohen also isolated EGF from human urine (Cohen

and Carpenter, 1975), where originally it was described as urogastrone and subsequently determined to be equivalent to EGF. The physical properties of EGF include a low molecular weight of approximately 6,000 Daltons and an isoelectric point of 4.6. Its amino acid sequence has been determined, revealing a single polypeptide chain of 53 residues with three intramolecular disulfide bonds (Taylor et al., 1972). Generally, EGF elicits a mitogenic response in cultured cells, and the response has been characterized experimentally by increased cell density and increased macromolecular synthesis, including DNA, RNA and protein (Cohen and Stastny, 1968; Hoober and Cohen, 1967a; 1967b; Westermark, 1976). Additionally, biological responses to EGF, which have been measured in various cells, involve ion transport, glycolysis, membrane alterations, viral growth and production of specific molecules (i.e. human chorionic gonadotropin, prostaglandin, etc.) (Carpenter and Cohen, 1979).

NGF activity was described by Bueker in 1948 in a mouse sarcoma. NGF was subsequently discovered in mouse submaxillary gland, snake venom, chick embryo, rat granuloma and other tissues and cells (reviewed in Gospodarowicz and Moran, 1976). The purified polypeptide

was determined to be a dimer consisting of chains of 118 amino acids bonded by three disulfide linkages (Angeletti and Bradshaw, 1971). Structurally and functionally, NGF is similar to both insulin and EGF, although it does not exhibit a mitogenic effect on cultured cells. NGF must be present in order to stabilize embryonic nerve cells in vitro, without NGF most nerve cells would disintegrate in culture (Bradshaw, 1978). Levi-Montalcini and Booker (1960a) demonstrated that NGF was necessary for cultured neurons to extend neurites. They also showed that the sympathetic nervous system of neonatal animals injected with anti-NGF antibodies was destroyed (1960b).

Another growth factor, which proved to be extremely interesting, was isolated from virally-transformed cells in culture and was, therefore, called transforming growth factor- β (TGF- β). TGF- β was shown to be a polypeptide of 25,000 Dalton molecular weight composed of two 12,500 Dalton subunits linked by disulfide bonds. TGF- β , although larger than EGF, was categorized with EGF due to similarites in structure (James and Bradshaw, 1984). TGF- β elicited opposing biological responses from various cells *in vitro*. Generally, fibroblasts were stimulated mitogenically, while epithelial cell lines appeared to be inhibited in proliferation. This contradictory activity

has not been fully explained through experimentation to date, but the resolution of this paradox may uncover some important information related to the positive and/or negative regulation of cell growth.

These various growth factors have been proposed to exert their influence by one of three modes of transmission (James and Bradshaw, 1984). The first model was related to normal hormone activity, or endocrine secretion. Endocrine secretion described the process whereby a molecule was produced by a cell or tissue, released into a transport system (i.e. the bloodstream), and subsequently, received by another cell far-removed from the original secreting cell. Alternatively, the released molecule might not have traversed such distances, but merely migrated to a neighboring, nonsecreting cell where it was received. Factors confined to a limited area illustrated the paracrine secretion model. Independent of the reaction of distant or neighboring cells, an actively secreting cell could contain receptors for the released molecules, thereby causing an effect in the manufacturing cell itself. This process has been referred to as autocrine secretion. Most of the growth factors have been demonstrated to operate in a paracrine fashion, but some that utilized

endocrine (insulin-like growth factors) or autocrine (TGFs) signaling have also been identified.

Regardless of the route of transport, these positive growth regulators induce mitogenesis by binding to specific receptors on the cell surface. The growth factors do not necessarily need to be internalized into the cell in order to initiate a response. Once a receptor-ligand complex is formed, a cascade of events proceeds within the cell. Among the intracellular changes that occur following receptor binding are influx of sodium ions, efflux of hydrogen ions, release of stored calcium ions, increased turnover of phosphatidyl inositols, and activation of protein kinase C (Deuel, 1987).

The multitude of changes that occur within the cell prior to division seem to be intricately connected. For example, the hydrolysis of phosphatidyl inositols produces diacylglycerol and inositol phosphates.

Inositol triphosphate may be a second messenger which couples membrane receptor binding to the release of intracellular calcium. At the same time, diacylglycerol triggers the activation of protein kinase C.

Molecules which may induce the hydrolysis of phosphitidyl inositols by acitivating phospholipase C are

what are termed G proteins. G proteins are membrane molecules composed of three polypeptide subunits which dissociate upon activation. The α -subunit of G proteins binds GTP, and in this state, is active. The active subunit of various G proteins not only may activate phosopholipase C, but also may stimulate or inhibit adenylate cyclase, which alters cellular cyclic AMP levels. Cyclic AMP triggers intracellular responses to hormones and possibly other regulatory factors.

The various positive growth regulators cause cell division via different pathways. Because the processes involved in mitosis are intertwined, it is plausible that a single inhibitory element could block the action of more than one stimulatory factor.

Negative Growth Regulators

I. Soluble Factors Responsible for Inhibition

The discovery of growth stimulatory factors in serum and various tissues, and the demonstration that cells in culture were density-dependent growth-arrested led some investigators to believe that antagonistic, inhibitory factors might also exist. Some researchers believed that such inhibitory factors might be released from cells in response to cell-cell contact and be present in conditioned culture media.

In 1974, the first cell-elaborated inhibitory factor was isolated by Lipkin and Knecht (1974). This diffusible factor which was isolated from culture media of a contact-inhibited, amelanotic melanoma cell line was capable of restoring contact inhibition to highly malignant melanocytes in vitro. The factor was demonstrated to contain protein and be of high molecular weight, but was not purified to homogeneity. Similarly, Holley et sl. (1978) have shown that the density-dependent regulation of growth of BSC-1 cells (African green monkey kidney) resulted, in part, from the accumulation of inhibitors in the medium. In 1980, Holley et sl. (1980) partially purified two high

molecular weight growth inhibitors from culture medium of BSC-1 cells. These factors were shown to inhibit thymidine incorporation in BSC-1 cells by 85%. The actual number of cells in culture was also limited by the inhibitory proteins. The action of these inhibitors could be counteracted by the addition of EGF. One of the BSC-1 cell inhibitors was later shown to be structurally similar to TGF-\$ (Tucker et al., 1984). In the last decade, several other groups have isolated and/or partially purified inhibitors from various sources. The hepatic proliferation inhibitor (HPI) was separated from a growth stimulatory fraction of rat liver homogenates by McMahon, Farrelly, and Type (1982). HPI was found to have a molecular weight of 26,000 Daltons and an isoelectric point of 4.65: its inhibitory activity was demonstrated by significantly reducing the number of colonies formed by Fischer nonmalignant rat liver cells.

Harel, Blat, and Chatelain (1985) isolated both inhibitory and stimulatory factors from 3T3 cell conditioned medium. The separated factors were IDF45, a 45,000 Dalton inhibitory polypeptide, and SDF10, a 10,000 Dalton stimulatory protein. IDF45 was demonstrated to inhibit the stimulation, by SDF10, of nucleic acid synthesis in 3T3 cells. Also from 3T3 cell conditioned

medium, Steck et al. (1982) isolated a polypeptide enriched fraction which they dubbed FGR-s (Fibroblast Growth Regulator-secreted). Partially purified, FGR-s contained two major polypeptides of molecular weights of 10,000 and 13,000 Daltons. This extract was shown to reduce the number of 3T3 cells per colony by two-thirds. Later, Hsu, Barry, and Wang (1984) produced a monoclonal antibody to FGR-s that bound the 13,000 Dalton protein and successfully neutralized the inhibitory activity.

In 1984, Boehmer et al. (1984) isolated and subsequently purified MDGI (Mammary Derived Growth Inhibitor) from normal bovine mammary glands. This protein had a molecular weight of 13,000 Daltons and inhibited the number of Ehrlich ascites mammary carcinoma cells in vitro by up to 40%. The MDGI inhibitory effect was not only abolished in culture by both EGF and insulin, but also neutralized by anti-inhibitor antiserum raised in mice. Interestingly, in 1987, Boehmer et al. (1987) discovered that antibodies prepared against MDGI cross-reacted with FGR-s (13 kD). There may be additional revelations such as that of the MDGI and FGR-s investigators. Such released inhibitors may function in an autocrine, paracrine and/or endocrine fashion, and presumably, these molecules usually act in an autocrine

or paracrine manner. As discussed above, a number of the inhibitors exert their effects on cells identical to those from which they were isolated. Logically, autocrine or paracrine secretion would explain density-dependent behavior of cells in culture, because cells within a confined space seem to signal themselves and/or each other to cease proliferation. Additionally, although positive growth regulators (growth stimulatory factors) have been shown to be present in serum, none of the inhibitors mentioned here have been isolated from serum.

II. Cell-associated Factors Responsible for Inhibition

Dulbecco and Stoker (1970) proposed that the cell growth inhibition of normal cells in vitro might be a result of cell-to-cell contact either directly or indirectly. As cell membranes approach one another, the receptors for various growth factors could be blocked, thus indirectly stopping cell proliferation. Later, Dulbecco (1970) showed that most likely "topoinhibition" was not caused by occlusion of receptors, but resulted directly from cellular interactions.

When Dulbecco showed that cell growth inhibition seemed to be a result of direct contact, other groups

(Burger, 1970; Sefton and Rubin, 1970) demonstrated that cells in culture lost contact inhibition when treated with proteolytic enzymes. Shortly thereafter, Pardee (1971) noted that during the process of cell division, the cell surface changed. Taken together, these observations suggested that some cell surface protein(s) which was(were) responsible for conferring density-dependent inhibition of growth to cells might indeed exist.

Following this lead, several groups attempted to isolate such a molecule. In 1976, Fisher and Koch (1976) isolated inhibitory peptides by treating HeLa cells with dilute pronase. HeLa cell surface peptides (HSP) were shown to inhibit protein synthesis of HeLa cells in culture and also in cell-free translation systems. HSP were incorporated into HeLa cells rapidly and appeared to associate with monosomes and polysomes. Using a similar method, Yaoi (1984) isolated inhibitory glycopeptides from primary chick embryo fibroblasts. These molecules reduced cell growth rate and thymidine uptake of chick embryo fibroblasts in vitro, but this inhibitory effect was partially destroyed by neuraminidase treatment and completely destroyed by periodate oxidation of the

sialoglycopeptides, suggesting that carbohydrates were responsible for the inhibition.

Finally, the Johnson laboratory isolated and purified glycopeptides from both mouse and bovine brain (see table for comparisons). The first glycopeptide inhibitors were proteolytically removed from mouse cerebral cortex cells in 1979 (Kinders, Johnson, and Rachmeler, 1979). These molecules, called BCSG (brain cell surface glycopeptides), had molecular weights of 10,000-25,000 Daltons and were shown to inhibit protein synthesis of various cells in culture. Subsequently, Kinders and Johnson (1982) isolated similar glycopeptides from boyine cerebral cortices. The first boyine derivative, which bound to Ulex europaeus agglutinin, thereby suggesting the presence of fucose, had an apparent molecular weight of 18,000 Daltons. The glycopeptide was demonstrated to inhibit cell proliferation and protein synthesis of BHK-21 (baby hamster kidney) cells in vitro. More recently, another bovine cerebral cortex glycopeptide has been purified (Sharifi et al., 1986) which also had a molecular weight of approximately 18,000 Daltons, but bound to Limulus polyphemus agglutinin and not to Ulex europaeus agglutinin, suggesting the presence of sialic acid

instead of fucose. The sialoglycopeptide (SGP) inhibitor reduced macromolecular synthesis and cellular mitosis in a wide spectrum of cell types (Fattaey, Johnson, and Chou, submitted). Binding of the SGP to a specific receptor (Bascom, Sharifi, and Johnson, 1986) was sufficient to produce the effects (Sharifi, Bascom, and Johnson, 1986), which were determined to be reversible. The inhibitor was capable of antagonizing the stimulatory effects of bombesin (Sharifi, unpublished), TPA (12-0-tetradecanoylphorbol-13-acetate) (Chou et al., 1987), and EGF (Bascom, Sharifi, and Johnson, 1987).

If these molecules were indeed cell surface proteins, then the groups highlighted above only isolated a portion of the inhibitory proteins by limited proteolysis. Concurrently, other investigators were working with cell membranes. Natraj and Datta (1978) extracted FGFR (Fibroblast Growth Regulatory Factor) from 3T3 cell membranes with urea. Previously, Whittenberger and Glaser (1977) had demonstrated that isolated 3T3 cell membranes were capable of inhibiting growth of cells in culture. Also in 1978, Whittenberger et al. (1978) proceeded to extract the inhibitory molecule from membrane fractions with the detergent, octylglucoside. Proteins were then partially purified from this

octylglucoside extract by Raben, Lieberman, and Glaser (1981).

TABLE. The information illustrated was collated from the Johnson laboratory. In each case, the cerebral cortices were prepared similarly for isolation of the various inhibitors. Briefly, cerebral cortex cells were treated with dilute protease, the elaborated proteins were ethanol precipitated and then extracted with chloroform-methanol.

Abbreviations are as follows: UEA-1=Ulex europaeus agglutinin, WGA=Wheat germ agglutinin, S-E HPLC=size-exclusion HPLC (high performance liquid chromatography), HI HPLC=hydrophobic interaction HPLC, HA HPLC= hydroxy-apatite HPLC, LFA=Limulus polyphemus agglutinin, ND=not determined, Met=methionine, Thy=thymidine, and Urd=uridine. Molecular weights in parentheses were determined using BioGel, others were determined by electrophoresis.

Comparison of inhibitors from cerebral cortices.

| Factor | Murine ¹ | I. Bovine2 | II. Bovine3 |
|--|--|---|--|
| Purification Steps | BiogelP-1001 UEA-11 | BioGelP-1002 UEA-12 CM agarose7 | DEAE agarose ³ WGA ³ S-E HPLC ³ HI HPLC ³ HA HPLC ⁹ |
| Molecular Weight | (26,000) ¹ (10,000) ¹ | (45,000) ² (10,000) ² 18,000 ² | 18,0003 |
| Isoelectric Point | ND | 8.1 ⁸ | 3.03 |
| Inhibition: | | | |
| Met incorporation Thy incorporation Urd incorporation Cell division | +1 ND -4 +4 | +2 ND ND +2 | +3 +10 +11 +10,12 |
| Cell cycle block | G25 | ND | G112 |
| Cell Specificity | A | В | С |

- A Primary cell lines most sensitive, transformed cell lines least sensitive, tumor cells sensitized by gangliosides. 1.6
- B Established cell lines and cells treated with
- gangliosides sensitive, transformed cells insensitive.2.7 $^{\circ}$ All cells tested to date sensitive, including primary,
- established, transformed, avian, and insect cell lines. 12
- 1 Kinders, Johnson, and Rachmeler, 1979.
- ² Kinders and Johnson, 1982.
- 3 Sharifi et al., 1986.
- 4 Kinders et al., 1980.
- 5 Kinders and Johnson, 1981.
- 6 Kinders, Rintoul, and Johnson, 1982.
- 7 Bascom et al., 1985.
- 8 Johnson, Kinders, and Sharifi, 1985.
- 8 Bascom et al., 1986.
- 10 Chou et al., 1986.
- 11 Bascom et al., in preparation.
- 12 Fattaey, Johnson, and Chou, submitted.

SECTION III
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SECTION IV

MANUSCRIPT: IN SEARCH OF--A MEMBRANE SIALOGLYCOPROTEIN
CELL GROWTH INHIBITOR

INTRODUCTION

The regulation of cell growth has been the subject of many studies and several investigators have isolated molecules from cells which inhibit cellular metabolism. A few of these inhibitory factors have been demonstrated to be membrane components, but none have been completely purified as membrane proteins. The possible residence of such molecules on the cell surface suggests that they may play an important role in cell-cell recognition, cell-cell communication and thus, the density-dependent growth behavior of cells in culture. The function of such growth inhibitors has been studied *in vitro*, using purified or partially purified molecules.

Sharifi et al. (1986) have isolated and purified, to near homogeneity, a glycopeptide inhibitor from intact bovine cerebral cortex cells that had an apparent molecular weight of 18,000 Daltons, an isoelectric point of 3.0 and has been shown to contain sialic acid by selectively binding to a Limulus polyphemus lectin affinity column.

The sialoglycopeptide (SGP) was purified by treating bovine cerebral cortex cells in suspension with dilute promase, followed by ethanol precipitation,

chloroform/methanol extraction, DEAE-agarose chromatography, wheat germ agglutinin chromatography, size exclusion HPLC, and hydroxyapatite HPLC (Sharifi et al., 1986).

The SGP inhibited the incorporation of radiolabeled methionine into cellular proteins and radiolabeled thymidine and uridine into cellular nucleic acids (Sharifi et al., 1986; Bascom et al., in preparation; Fattaey, Johnson, and Chou, submitted). The lack of incorporation of such precursors into cellular products suggested an inhibition of cell division. It has also been shown that the SGP reduced the proliferation of cells, as determined by direct cell count (Chou et al., 1987; Fattaey, Johnson, and Chou, submitted). The inhibitor appeared to synchronize 3T3 cells in the G1 phase of the cell cycle and upon removal of the SGP, the cells resumed proliferation.

In order to supress cell proliferation, the SGP did not need to be internalized by the cells. Binding to a specific cell surface receptor was sufficient for the SGP to mediate its inhibition (Bascom, Sharifi, and Johnson, 1986; Sharifi, Bascom, and Johnson, 1986). Additionally, the inhibitor was shown to be a potent antagonist of the mitogenic activity of growth factors and a tumor

promoter. The SGP successfully inhibited the stimulation of cells by bombesin (Sharifi, unpublished), epidermal growth factor (Bascom, Sharifi, and Johnson, 1987), and TPA (12-0-tetradecanoylphorbol-13-acetate) (Chou et al., 1987).

As stated previously, the SGP was isolated by proteolytic cleavage from the surface of bovine cerebral cortex cells, thereby presumably generating a fragment of a resident membrane glycoprotein, or "parental" glycoprotein. The inhibitory fragment was soluble in water and therefore, unlike the typical hydrophobic proteins found in membranes, was convenient to purify by routine biochemical procedures that employ aqueous solvents.

Prior to this study, rabbit polyclonal anti-SGP antibody was produced and the availability of this antibody provided an opportunity to assess a "one-step" purification scheme. Theoretically, the cell growth inhibitor (SGP) and/or its "parental" membrane protein could be isolated by applying an extract from cerebral cortex cells to a matrix coupled with anti-SGP polyclonal antibody. Such a strategy was employed in the following attempt to purify the "parental" sialoglycoprotein in its native state, as a membrane molecule.

MATERIALS AND METHODS

Isolation

Slices of bovine cerebral cortex tissue were removed from fresh bovine brain, placed in ice-cold DMEM (Dulbecco's Modified Eagle Medium, GIBCO Laboratories Life Technologies, Inc., Grand Island, NY) containing 25 mM HEPES buffer (pH 7.1) and weighed. A cell suspension was prepared by repeated pipetting. The cells were pelleted by centrifugation at 3,000 rpm for 5 minutes and resuspended in 1-2 ml ice-cold solubilization buffer per gram of wet tissue. Solubilization buffer contained 1% w/v octyl-β-D-glucopyranoside (Behring Diagnostics. La Jolla, CA), 10 nM phenylmethylsulfonylfluoride (Serva Fine Biochemicals, Inc., Westbury, NY), 5 nM phosphoramidon (Sigma Chemical Company, St. Louis, MO), 10 nM leupeptin (Serva Fine Biochemicals, Inc., Westbury, NY), 10 nM pepstatin A (Sigma Chemical Company, St. Louis, MO), in phosphate buffered saline (PBS), where PRS was composed of 0.9% w/v sodium chloride, 10 mM sodium phosphate, pH 7.2. After incubation on ice for 30 minutes with periodic mixing, the suspension was centrifuged at 5,000 rpm for 5 minutes. The supernatant

fluid was collected and re-centrifuged at 100,000 x g for one hour. The supernatant liquid was then concentrated to a final volume of 10 ml by ultrafiltration using a Diaflo YM100 membrane (Amicon Corporation, Danvers, MA).

Protein Determination

Protein content was determined by the BCA (bicinchoninic acid) protein assay (Pierce Chemical Company, Rockland, IL) using bovine serum albumin in PBS containing 1% (w/v) octyl- β -D-glucopyranoside (octylglucoside) as a standard.

Polyclonal Antibody Affinity Column

Typically, 1.5-2 ml activated agarose slurry, Affigel 10 (BioRad Laboratories, Richmond, CA), were washed with at least 5 column volumes of ice-cold distilled water followed by approximately 5 column volumes of PBS. Approximately 200 µg of polyclonal antibody, adjusted to a final volume of 1 ml, were added to the column matrix and rotated on a Fisher Hematology/Chemistry Mixer (Fisher Scientific, Pittsburgh, PA) continuously, overnight, at 4°C. The Affi-gel 10 was then washed with at

least 10 column volumes of PBS and fractions were saved to determine the amount of uncoupled antibody. Unreacted Affi-gel 10 was then blocked by washing successively with several volumes of 3 M ammonium thiocyanate, 50 mM ethanolamine (pH 11.0) and/or 2 M glycine (pH 2.5). Excess blocking agents were then removed by extensive washing with 100-500 column volumes of PBS, followed by equilibration with 10 column volumes of PBS containing 1% (w/v) octylglucoside.

Polyclonal Antibody Affinity Chromatography

Concentrated, solubilized bovine cerebral cortex proteins were applied to the polyclonal antibody affinity column and incubated at 4°C, overnight. The column was washed with PBS containing 1% (w/v) octylglucoside until no absorbance at 280 nm was detectable. Elution buffer (1% w/v octylglucoside in either 50 mM ethanolamine or 2 M glycine) was added to elute antibody-bound proteins. Samples were collected until absorbance was zero. A single protein peak was eluted and the collected fractions were pooled, dialyzed against dilute PBS containing 1% (w/v) octylglucoside, dried in a Speed Vac

Concentrator (Savant Instruments, Inc., Hicksville, NY) and stored at -20°C.

SDS-PAGE

Proteins were diluted 1:2 (v/v) in sample buffer (36 mM sodium phosphate, pH 7.2 with 35% glycerol, 4% sodium dodecyl sulfate, 0.1% bromphenol blue with or without 18% β -mercaptoethanol). Samples were incubated at 100°C on a heating block for 5-10 minutes and then were electrophoresed in a 5%-20% polyacrylamide gradient, essentially by the method of Laemmli (1970). Molecular weight standards (prestained, Bethesda Research Laboratories Life Technologies, Inc., Gaithersburg, MD) were myosin H-chain (200,000), phosphorylase B (97,400), bovine serum albumin (68,000), ovalbumin (43,000), α -chymotrypsinogen (25,700), β -lactoglobulin (18,400) and lysozyme (14,300).

Isoelectric Focusing

Isoelectric focusing was performed by a technique similar to the method of Vesterberg (1972). The gel was poured between a Gel-Bond (FMC Corporation, Rockland, ME) and glass plate sandwich, using the following solution: 5% (v/v) acrylamide stock, 10% (v/v) glycerol, 0.9% (v/v) Triton-X 100 (CalBiochem, La Jolla, CA), 0.3% (v/v) Ampholine pH 4-6 (LKB Wallac, Finland), 0.3% (v/v) Ampholine pH 5-7, 0.6% (v/v) Ampholine 9-11, 4.6% (v/v) Ampholine pH 3.5-10, and 0.05% (w/v) ammonium persulfate. The electrodes were saturated with 0.5 N sodium hydroxide and 0.5 N acetic acid. After focusing for 90 minutes at 25 Watts, the gel was removed. The pH gradient was measured using a flat-bed electrode, unless isoelectric focusing standards were used. Broad Range IEF Standards from BioRad included phycocyanin-pI 4.65, β-lactoglobulin B-pI 5.10, bovine carbonic anhydrase-pI 6.00, human carbonic anhydrase-pI 6.50, equine myoglobin-pI 7.00, whale myoglobin-pI 8.05, chymotrypsin-pI 8.80, and cytochrome c-pI 9.60.

Size Exclusion HPLC

A molecular sieving column (TSK G3000, Phenomenex, Rancho Palos Verdes, CA) was equilibrated with PBS containing 1% (w/v) octylglucoside, samples were applied, and 1 ml aliquots collected, at a flow rate of 1 ml/minute. Absorbance was monitored at 280 nm.

Limulus polyphemus Agglutinin (LPA) Chromatography

About 2 ml (final column volume) of LPA (E-Y Laboratories, San Mateo, CA) were equilibrated with at least 10 column volumes of 10 mM Tris buffer, pH 8.0, containing 10 mM calcium chloride and 1% (w/v) octylglucoside. Bovine cerebral cortex solubilized proteins at various stages of purification were added to the matrix and rotated on a Fisher Hematology/Chemistry Mixer (Fisher Scientific, Pittsburgh, PA) continuously, at 4°C, overnight. The LPA was washed until no absorbance at 280 nm was detectable. Subsequently, lectin-bound molecules were eluted with 100 mM sialic acid, pH 8.0 with 0.1% EDTA and 1% (w/v) octylglucoside. Eluted fractions were dialyzed against dilute PBS containing 1% (w/v) octylglucoside and dried in a Speed Vac Concentrator (Savant Instruments, Inc., Hicksville, NY).

Reduction and Blocking

Bovine cerebral cortex protein extracts were reduced by adding β -mercaptoethanol to a final concentration of 9% (v/v) and incubating at room temperature for 2 hours. The sulfhydryl groups were alkylated by dialysis against 100 mM iodoacetamide for 4 hours followed by dilute (0.1 mM) iodoacetamide at 4°C, overnight.

Cell Culture

Swiss 3T3 cells were maintained as monolayer cultures in humidified, 37°C incubators containing 95% air and 5% carbon dioxide, as described by Kinders et al., (1979). Cells were grown in Dulbecco's Modified Eagle Medium (GIBCO Laboratories life Technologies, Inc., Grand Island, NY), supplemented with 10% calf serum (Hazelton Dutchland, Inc., Denver, PA) and were harvested for assay by scraping culture flasks with sterile rubber policemen.

Biological Inhibitory Activity

The method utilized to determine the capability of a molecule to inhibit protein synthesis of cells has been described by Kinders et al., (1979). Basically, Swiss 3T3 cells were harvested from subconfluent monolayers after washing briefly with methionine-free Minimal Essential Medium containing 25 mM HEPES buffer (MEM-

HEPES). The cell suspension was then centrifuged; the pelleted cells were resuspended and counted in a hemacytometer, and diluted to provide 2-5 x 10^5 cells in 100 µl of warm (37°C), methionine-free MEM-HEPES. Potential inhibitory substances were pipetted into 13 X 100 mm test tubes at different concentrations, each in a final volume of 40 µl in PBS. Cell aliquots were then added and the tubes were incubated, at 37°C, for 30 minutes. At the end of this incubation, 2.0 µCi of 35Smethionine in 10 µl of methionine-free MEM were added and the tubes immediately reincubated, at 37°C, for 10 minutes. Incorporation was stopped by the addition of 2 ml ice-cold HKM buffer (10 mM HEPES, 120 mM potassium chloride, and 5 mM magnesium chloride, pH 7.1) and centrifugation of the cells, at 2,000 rpm, for 3 minutes. After removal of supernatant liquid, the cells were lysed with distilled, deionized water and 1 N sodium hydroxide, followed by precipitation of proteins with an equal volume of ice-cold 20% trichloroacetic acid. Precipitated molecules were collected by centrifugation and the previous steps repeated. On the third repetition, trichloroacetic acid was omitted and a 100 μ l aliquot was removed for liquid scintillation counting.

Immunoblotting

Samples were either directly spotted onto pre-wet nitrocellulose (Schleicher and Schuell, Inc., Keene, NH) using a Minifold (Schleicher and Schuell, Inc., Keene, NH) or electrophoresed and electroblotted using a Transphor Electrophoresis Cell (Hoefer Scientific Instruments, San Francisco, CA) by the method of Towbin, et al., (1979). Following transfer, the nitrocellulose membrane was dried and then blocked overnight, at room temperature, in blocking buffer (0.1 mM EDTA, 0.01% w/v sodium iodide, 5 mM sodium chloride, 1 mM Tris, pH 7.0 and 0.3% Tween 20). The membrane was subsequently washed 5 times, for 5 minutes each, with wash buffer (1% w/v sodium chloride, 100 mM Tris, pH 7.5 and 0.3% Tween 20). Antibody was added, in blocking buffer, and the membranes were incubated overnight, at room temperature. The blot was washed again, followed by the addition of 125Iradiolabeled staphylococcal Protein A at 100,000 CPM per ml of blocking buffer. Finally, the membrane was washed. dried and used to expose an X-ray film (Agfa-Gevaert, Belgium).

RESULTS

Isolation

The strategy designed to purify the parental sialoglycoprotein inhibitor is illustrated in Figure 1. Fifteen independent isolates from fresh bovine cerebral cortex tissue cell suspensions were studied. The cells were pelleted by low speed centrifugation and extracted with 1% (w/v) octylglucoside. Only 4.0% of the total cellular protein was extracted by this method. Following solubilization, the cells were again centrifuged; the supernatant fluid, containing detergent-extracted proteins, was retained and concentrated by ultrafiltration. The concentrated extract contained 3.0% of the total cerebral cortex cell protein, as measured by BCA protein assay (Table 1).

Polyclonal Antibody Affinity Purification

The detergent-extracted proteins were applied to a polyclonal antibody affinity column and subsequently eluted with either ethanolamine or glycine, containing 1% (w/v) octylglucoside (Fig. 2). In either case, the

collected fractions were immediately neutralized, dialyzed against dilute PBS, and analyzed by electrophoresis. In parallel experiments with the 18,000 Dalton SGP inhibitor, it was determined that the sialoglycopeptide was not efficiently released from a polyclonal antibody affinity column by 50 mM ethanolamine. Subsequently, biologically active SGP was successfully eluted from the antibody column with less than ten column volumes of 2M glycine, pH 2.5.

Because of the previous observation, the majority of "parent" isolations were conducted using glycine as elution buffer. The amount of protein recovered by elution with 50 mM ethanolamine was 0.002% of the total cerebral cortex cell protein compared to 0.003% of the total cerebral cortex cell protein eluted with 2 M glycine, pH 2.5 (Table 1).

Molecular Analysis of Polyclonal Antibody-Eluted Material

Proteins bound to the polyclonal antibody affinity column were subjected to SDS-polyacrylamide gel electrophoresis. Non-reduced preparations failed to enter the separating gel, whereas, reduced preparations demonstrated the presence of four distinct protein bands

of apparent molecular weights of 175 kD, 135 kD, 66 kD, and 56 kD (Fig. 3). This indicated that the protein (or proteins) might be a large complex that included four subunits connected by disulfide bonds.

Although the non-reduced proteins failed to enter polyacrylamide gels, it was possible that these high molecular weight species might be discernable by size-exclusion HPLC (high performance liquid chromatography). Molecular sieving HPLC could have provided a convenient means of separation of the antibody eluted molecules, but the chromatographs from several preparations were inconsistent (Fig. 4). The hydrophobicity of these membrane proteins may have caused them to aggregate randomly into various conformations in different isolates. These complexes of diverse molecular size could have produced different elution times and peak areas in size-exclusion HPLC. In any event, it was impossible to assess the molecular weight species by HPLC.

Interestingly, the four reduced bands present in SDS gels were no longer obvious upon isoelectric focusing.

Isoelectric focusing of the polyclonal antibody affinity purified material, revealed only a slight difference between non-reduced and reduced samples. Focusing did

not produce distinct protein bands, but did show that the molecules present were acidic and heterogeneous with regard to their isoelectric points (Fig. 5). This heterogeneity suggested that the proteins were glycosylated and/or sialylated to varying degrees.

In order to determine if the antibody-bound molecule(s) contained N-acetylneuraminic acid (NANA), binding to a NANA-specific lectin, Limulus polyphemus agglutinin (LPA), was examined. It appeared that at least a portion of the isolated material contained sialic acid because, on average, forty percent of the protein applied to the LPA column bound, and in an early experiment neuraminidase-treated proteins were shown not to bind to the LPA. However, the amount of glycoprotein, in separate preparations, eluted from LPA by competition with sialic acid varied from three to seventy-six percent bound of the total protein applied. Polyacrylamide gels revealed similar staining patterns in both the LPA-bound and LPA-unbound fractions.

Reduction and Blocking of Polyclonal Antibody-Eluted

Due to the problem of aggregation of the molecules when under non-reducing conditions, it was thought that reduction followed by alkylation to prevent reformation of disulfide bonds might aid in the elucidation of the specific subunit directly related to the sialoglycopeptide inhibitor. By decreasing the probability for subunits to bind together in disulfide linkages, it seemed possible that the molecules might remain separate during analysis or that, perhaps, the subunit most similar to the SGP might be less hydrophobic than the other subunits.

Reduction was accomplished using a final concentration of 9% (v/v) β -mercaptoethanol and subsequent alkylation was performed with 100 mM iodoacetamide. Sufficient blocking was demonstrated to have occurred by subjecting the alkylated molecules to SDS-PAGE in the absence of reducing agents (Fig. 6). The proteins stained by Coomassie Blue were analogous to those seen previously under reduced conditions (Fig. 3).

Reduction and alkylation did not appear to facilitate separation and identification of the subunit directly related to the SGP. For example, a reduced and blocked preparation was applied to an LPA column, producing results which were similar to those seen with non-reduced, non-alkylated protein. Approximately fifty percent of the sample applied to the LPA column was bound and released. Which of the four bands seen in SDS-PAGE actually had NANA residues was not well-defined by electrophoretic analysis of the LPA-bound and unbound fractions. Based on SDS gels, all of the proposed subunits recovered from the lectin column contained sialic acid. Nearly identical staining patterns were observed in both bound and unbound samples.

Biological Activity of Polyclonal Antibody-Eluted Material

Although the molecular characterization was ambiguous, auspicious results were obtained from a biological activity assay. One of the initial criteria demanded that the "parent" to the sialoglycopeptide cell growth inhibitor should be capable of inhibiting cell

growth and/or protein metabolism. Polyclonal antibody purified material was dialyzed to remove detergent and tested for its ability to inhibit Swiss 3T3 cells from incorporating radiolabeled methionine.

In one such experiment, protein synthesis was effectively inhibited by the putative "parent" molecule. A dose-responsive inhibition of protein synthesis was observed in Swiss 3T3 cells treated with various concentrations of antibody-purified proteins (Fig. 7). Additional trials of the biological activity assay did not reproduce the inhibition described here; therefore, the conclusion that the isolated protein was the "parent" of the SGP inhibitor was not experimentally confirmed.

The amount of potential "parent" protein used in biological activity assays was based on experiments performed with the inhibitory peptide. In preliminary isolations of the "parent", the protein was demonstrated to have an apparent molecular weight of 180,000 Daltons, which was ten times as large as the peptide inhibitor. At the time this assay was performed, 2 µg of SGP (peptide) was equivalent to one unit of biological activity, which was defined as that amount of protein that inhibited methionine incorporation by 25%.

Consequently, 20 µg of the suspected "parent" protein was the minimum used in cell protein synthesis experiments.

After heating the antibody-isolated fraction to 100°C for 30 minutes, no inhibition of methionine incorporation was evident in Swiss 3T3 cells, suggesting that the inhibition described above resulted from the presence of a non-denatured protein.

The inhibition was not due to the presence of detergent in the protein samples, because the dialysate from the octylglucoside removal step was used as a control. Additionally, dilutions of detergent were assayed in a separate experiment which demonstrated that up to 0.02% octylglucoside would not significantly inhibit protein synthesis in Swiss 3T3 cells (Table 2).

Immunoblotting of Polyclonal Antibody-Eluted Material

Since at least one experiment indicated that a protein in the polyclonal antibody purified fraction was inhibitory, it seemed logical that the molecule responsible for inhibition could be identified by immunoblot analysis. Polyclonal antibody-isolated material, LPA-eluted material, and crude octylglucoside solubilized material were subjected to immunoblot

analysis. Blots were performed by direct application of samples to nitrocellulose (dot blots) and by SDS-PAGE followed by electrophoretic transfer to nitrocellulose (Western blots). Efficient transfer of proteins was demonstrated by negative gel staining with Coomassie Blue and positive membrane staining with Ponceau S. The 18,000 Dalton SGP inhibitor, which had been used as the immunogen, successfully bound polyclonal antibody in an immunoblot. During the course of this study, no reactive species from either affinity purified samples or crude octylglucoside extracted samples were detected in immunoblots. However, more recently generated antiserum does react with detergent solubilized and immunoaffinity column purified proteins in a dot blot.

Non-specific Binding to the Polyclonal Antibody Column

Based on the lack of evidence that the isolated proteins consistently reacted with polyclonal antibody, it was suspected that the immunoaffinity column may not have been binding specifically to SGP-related molecules. Non-specific binding was tested experimentally by applying solubilized material to an uncoupled, blocked agarose column. In this manner, it was shown that

detergent-extracted proteins actually bound to the uncoupled agarose to nearly the same extent that they bound to antibody-coupled agarose (Table 3). This suggested that the concentration of antibody ligated to the matrix was too low or that the isolated proteins had some affinity for the matrix itself.

In order to reduce such non-specific binding, a new immunoaffinity column was prepared using a 10-100 fold greater concentration of antibody and an uncoupled agarose column was inserted into the isolation strategy as a pre-column. Neither of these procedures appeared to facilitate the purification of an immunoreactive protein, as measured by blotting.

DISCUSSION

The ultimate goal of this project was to purify the native form, or "parent" of the bovine cerebral cortex cell surface sialoglycopeptide growth inhibitor. During the process of attempting to purify the membrane sialoglycoprotein, valuable information was discovered.

The purification of the native membrane sialoglycoprotein cell growth inhibitor proved to be a more formidable exercise than originally perceived.

Purification was difficult for several reasons. First, the "parent" sialoglycoprotein probably constituted a mere fraction of the total cellular proteins, because its derivative was a minor bovine brain protein. Only microgram quantities of the SGP cell growth inhibitor were separated from over 100 grams of cerebral cortex tissue (Sharifi et sl., 1986).

The isolation of some protein which was relatively abundant would have been a less arduous task. For example, the isolation of comparatively large quantities of hemoglobin would have been facilitated by its known high concentration in blood: 14-16 grams of hemoglobin per 100 ml of whole blood (from New England Journal of Medicine, 1980).

While hemoglobin is relatively abundant in blood, growth factors are present in much lower concentrations in serum and growth factor receptors are comparatively scarce on cell surfaces. One group of researchers was fortunate that a unique cell line was discovered. The recognition that an epidermoid carcinoma cell line, A431, specifically bound approximately ten-fold more EGF than other human cell lines (Fabricant, DeLarco, and Todaro, 1977) enhanced the purification of the EGF receptor (Cohen, Carpenter, and King, 1980). In the present study, an abnormally rich source of the desired molecule might have been utilized, if such a source were available or identifiable.

As in the isolation of the SGP, it was speculated that brain tissue, because of its growth-arrested state in vivo, was a propitious source of potential cell growth inhibitory elements. It was known, as stated above, that the 18,000 Dalton SGP inhibitor was present on the surface of bovine cerebral cortex tissue in limited quantities, but it was uncertain whether or not the number of parental molecules would correlate with the number of SGP molecules.

Isolating a minute portion of the total cellular protein was difficult enough, but the fact that the

extracted membrane proteins were hydrophobic and predisposed to aggregation added another dimension to the dilemma. Further increasing the magnitude of the purification problem was the finding that at least some of the solubilized proteins tended to bind non-specifically to the immunoaffinity column.

Some of these difficulties must have been encountered by others in the pursuit of membrane inhibitory molecules. Several researchers have claimed to purify such negative growth regulators, but have not demonstrated conclusively that their preparations were indeed pure.

The "purification" of a growth inhibitor from Ehrlich ascites mammary carcinoma cells by Boehmer et al., (1984) demonstrated the presence of an inhibitory protein of approximate molecular weight between 12,000 and 14,000 Daltons. The molecular weight of the major protein isolated by Grosse's group was later described as 13,000 Daltons (Boehmer et al., 1985), but the 1251-labeled inhibitory fraction appears as a single protein band only after fixing and washing gels as if preparing for silver staining. When gels were not prepared for silver staining, multiple bands were present. Biological activity was tested in samples which were not treated for

silver staining, therefore, inhibitory properties cannot be unconditionally attributed to the 13,000 Dalton protein. Because it was not clear that the major protein discussed above was responsible for cellular inhibition and, in an earlier publication (Boehmer et al., 1984), the greatest proportion of inhibitory activity was attributed to a fraction which displayed an almost invisible protein band, the possibility exists that a minor protein actually affected cell growth.

Another group that happened upon difficulties in working with inhibitory molecules was Luis Glaser's lab. Glaser and coworkers isolated 3T3 cell membranes which were shown to inhibit cell growth (Whittenberger and Glaser, 1977). The published work was criticized because Glaser et al. had not purified the specific inhibitory element from the cell membrane. The arguments against Glaser included the fact that the membrane fragments could, in essence, be "smothering" the cells by not allowing interaction of growth factors in the medium with receptors on the cell surface. To refute the contention that cells treated with membrane fragments were inhibited due to lack of stimulation by resident growth factors, Whittenberger and Glaser (1978) submitted an elegant

experiment which displayed that limited diffusion was not responsible for interruption of cell growth in culture.

Whittenberger et al., (1978) subsequently solubilized the membrane fragments and demonstrated that the extracted proteins could also inhibit cell metabolism. Three years later, the growth inhibitory proteins remained "partially purified" (Raben, Lieberman, and Glaser, 1981), suggesting that the researchers had not been successful at separating a homogeneous protein.

One of the biggest obstacles in the present study was the tendency of the extracted molecules to aggregate. There is a precedent for such difficulties: Patt and Houck (1980) reported that "chalones", endogenous mitotic inhibitors, had an "aggregative avidity...for various macromolecules."

The propensity for the isolated proteins to aggregate may have been the cause of some of the inconsistent experimental results. The inconsistencies may have resulted from variations in the arrangement of the aggregates. For instance, one of five biological activity assays was successful in demonstrating inhibition of cellular protein synthesis. This may have been because, in that particular trial, the protein complex was arranged such that the inhibitory portion of

the "parent" molecule was available for binding to the surface of 3T3 cells, thereby signaling the cells to cease proliferation.

Aside from its relatively low abundance on cerebral cortex cells and the aggregative behavior of the native sialoglycoprotein, a substantial inconvenience was the preparation and utilization of an immunoaffinity column. This project was based on the supposition that a "parent" molecule actually existed on the surface of bovine cerebral cortex cells. Furthermore, it was speculated that the membrane glycoprotein in its native conformation contained an epitope which would bind to and was available to polyclonal antibodies raised against the SGP inhibitor.

Consequently, the preparation of an immunoaffinity column using the previously isolated polyclonal antibodies was presumed to be of vital significance.

While utilizing the antibody affinity column, nonspecific binding of the solubilized proteins to the column became apparent. The interpretation of this result was that either the extracted molecules had a high affinity for the column (matrix, etc.) or the antibodies had a low (or no) affinity for the "parent" glycoprotein. The antibody may not have bound the "parent" molecule

because the membrane glycoprotein complex conformation was such that the epitope(s) was (were) hidden. More than likely, a combination of the former was responsible for these results.

The polyclonal antibody used in the present study had been passed through an affinity column coupled with purified SGP (Sharifi, unpublished). The affinity-purified antibody was shown to have high affinity for the peptide inhibitor in radioimmunoassays (Fleenor, unpublished), but assays with tissue culture cells and bovine cerebral cortex cells failed to display immunofluorescence (Fleenor, unpublished). The inability to demonstrate fluorescence was attributed to a cell surface concentration of SGP that was under the limits of detection of the assay. Alternatively, the epitope(s) might have been masked by other cell surface proteins or the native protein itself.

A potential solution to the dilemma of non-specific binding to the immunoaffinity column would be to increase the quantity of antibody coupled to the column matrix, provided that the non-specific binding described here was due to proteins binding to the matrix and/or the plastic column. The evidence presented by the binding of molecules to the uncoupled agarose strongly suggests that

this could be the case. It is plausible that the purification of the parental sialoglycoprotein could be enhanced by the availability of a greater supply of antibody and/or by a higher affinity antibody.

The affinity columns prepared in this study were not coupled to maximum capacity, which may have provided extra matrix surface area to which the proteins might adhere. Logically, if the agarose beads were coupled to capacity, then there would be fewer potential binding sites on the matrix itself. Other laboratories have encountered difficulties in protein purification on activated agarose columns which were coupled at low densities (J. Hughes, personal communication).

Additionally, antibody with a greater affinity or alternative binding site might alleviate some of the problems mentioned. The study described here may have resulted in the isolation of the native membrane sialoglycoprotein, but because of the difficulties encountered, the "parent" molecule has not been conclusively identified. Newly generated antibody could conceivably react positively in an immunoblot and, thereby, remove some of the ambiguity and doubt described above. Such an antibody appears to have been produced recently. This novel antiserum has not only bound crude

bovine cerebral cortex protein, detergent-extracted protein, and several of the original brain isolates in a dot blot, but also bound potential "parent" molecule to a freshly prepared immunoaffinity column. Thereafter, an immunoreactive species was identified in the eluate from the new column. These preliminary results indicate that an antigenic determinant is actually present and accessible in the octylglucoside solubilized material. Hopefully, this antibody will assist in the search for the native membrane sialoglycoprotein, provided the other obstacles described here are also conquered.

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TABLE 1. Purification of the parental sialoglycoprotein.

| Purification Step | Protein (mg/g wet tissue) | Total (%) |
|--|---------------------------|-----------|
| Cerebral Cortex Cells | 125.0 | 100 |
| Octylglucoside Extract | 5.5 | 4 |
| Concentrated Extract | 3.9 | 3 |
| Polyclonal Antibody Bound (released with glycine) | 0.004 | 0.003 |

Protein concentrations were determined by the BCA protein assay as described in Materials and Methods.

Data shown are from a typical bovine cerebral cortex preparation.

TABLE 2. Inhibition of methionine incorporation by octylglucoside.

| Octylglucoside Concentration (% w/v) | Inhibition (% of control) |
|--------------------------------------|---------------------------|
| 0 (control) | 0 |
| 0.012 | 0 |
| 0.025 | 6 |
| 0.05 | 12 |
| 0.1 | 40 |
| 0.25 | 99 |
| 0.5 | 99 |
| 1 0 | 99 |

Alignots of 100 μ l methionine-free MEM-HEPES containing 2-5 x 10⁵ Swiss 3T3 cells were incubated with 40 μ l samples of various concentrations of octylglucoside in PBS for 30 minutes, at 37°C. At the end of this incubation, the cells were radiolabeled with 2.0 μ Ci of 35S-methionine, at 37°C, for 10 minutes and processed as described in Materials and Methods. These data represent the averages of duplicate samples.

TABLE 3. Non-specific binding of proteins to uncoupled agarose.

| Column Type | Protein Bound (µg) | Total (%) |
|---|-----------------------|-----------|
| Polyclonal Antibody Coupled, Blocked Affi-gel 10 | 20 | 1.9 |
| Uncoupled, Blocked Affi-gel 10 | 18 | 1.7 |
| Agarose | 31 | 2.9 |

Aliquots of 1 mg of solubilized bovine cerebral cortex proteins were applied to the columns, incubated at 4°C, overnight, and subsequently eluted with 2 M glycine, pH 2.5. Protein concentrations were determined by BCA protein assay. These data represent the averages of three experiments.

Figure 1. General purification procedure for the isolation of the bovine cerebral cortex cell membrane sialoglycoprotein.

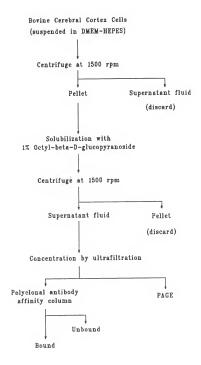
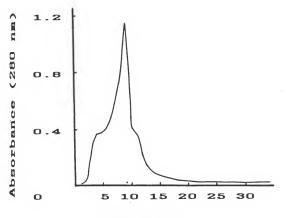


Figure 2. Polyclonal antibody affinity column elution of the potential "parent" sialoglycoprotein. The octylglucoside extracted fraction containing approximately 5 mg protein was loaded onto Affi-gel 10 previously coupled with polyclonal antibody. The single eluted peak was collected and analyzed.



Fraction

Figure 3. SDS-PAGE of fractions purified by immunoaffinity chromatography. Three separate preparations were electrophoresed in a gel with a linear gradient of 5-20% polyacrylamide. The preparations were analyzed under non-reducing (Lanes 1-3) and reducing (Lanes 5-7) conditions. Each lane was loaded with 5 µg protein. Lanes 1 and 5 contain samples from the same brain preparation, as do Lanes 2 and 6, and Lanes 3 and 7. Lane 4 contains molecular weight standards.

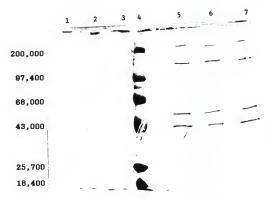
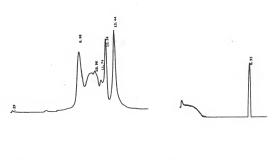


Figure 4. Size-exclusion HPLC analysis of immunoaffinity eluted fraction. Samples containing 10 µg of protein were loaded on a TSK G3000 HPLC column previously equilibrated with phosphate-buffered saline, pH 7.2. The attenuation was set at 16 and the flowrate was 1 ml per minute. Panels A-D illustrate HPLC profiles from four separate brain isolates. Elution times are as indicated.





; D

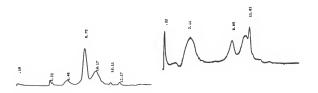


Figure 5. Isoelectric focusing of immunoaffinity eluted fraction. The purified protein was radioiodinated and isoelectric focused for 1.5 hours at 25 Watts. Samples of 100,000 CPM were loaded into each lane. Sample A was reduced while Sample B was not reduced.

pH A B
8.1

7.3

6.3

5.6

4.3

4.3

Figure 6. SDS-PAGE of reduced and alkylated immunoaffinity eluted fraction. A sample of 5 μ g of blocked protein was electrophoresed in a gel with a linear gradient of 5-20% polyacrylamide in the absence of reducing agents. Lane 1 is the blocked sample and Lane 2 contains molecular weight standards.

1 2

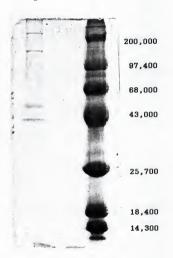
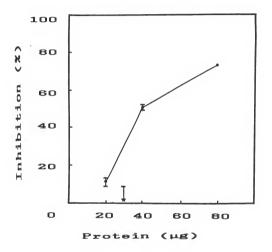


Figure 7. Inhibition of 3T3 cell protein synthesis by an immunoaffinity eluted fraction. Cells in 100 μl of DMEM-HEPES were incubated at 37°C in the presence of 20 μl of the isolated protein in PBS, as described in Materials and Methods. Assay was performed in duplicate and results are expressed as percentage of inhibition as compared with control cultures, which received only PBS. The vertical bars depict the range of duplicate samples and the * indicates a sample which was heated to 100°C for 30 minutes prior to addition to cells.



IN SEARCH OF: A MEMBRANE SIALOGLYCOPROTEIN CELL GROWTH INHIBITOR

bу

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B.S., Kansas State University, 1985

AN ABSTRACT OF A THESIS

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ABSTRACT

Density-dependent control of cell growth in culture may result from specific inhibitory molecules in cell membranes interacting with specific receptors in neighboring cell membranes. A molecule with such potential has been proteolytically cleaved from the surface of bovine cerebral cortex cells and has been shown to inhibit cell division, protein synthesis, DNA, and RNA synthesis of several cultured cell lines. present study utilized polyclonal antibody against the sialoglycopeptide inhibitor in an attempt to isolate the intact, native membrane protein from which the inhibitor was originally derived. Bovine cerebral cortex cells were solubilized with octyl-β-D-glucopyranoside (octylglucoside) and the solubilized molecules subjected to immunoaffinity chromatography. The polyclonal antibody bound fraction was examined by SDS-PAGE, IEF, HPLC, lectin affinity chromatography and immunoblot analysis. The isolated material was also tested for biological activity in a protein synthesis inhibition assay. Upon reduction in an SDS polyacrylamide gel, the antibody-bound fraction displayed four distinct molecular weight bands. The extracted protein inhibited protein synthesis in Swiss 3T3 cells; however, this inhibition

was not predictably repeatable. Immunoblots failed to reveal antibody binding activity in either the affinity purified material or in crude detergent solubilized fractions. Based on the evidence presented, it cannot be concluded that the fraction isolated was indeed the native membrane sialoglycoprotein. However, at the time of this writing, preliminary results indicate that octylglucoside extracted molecules react with more recently generated antiserum.